

Fig. 1

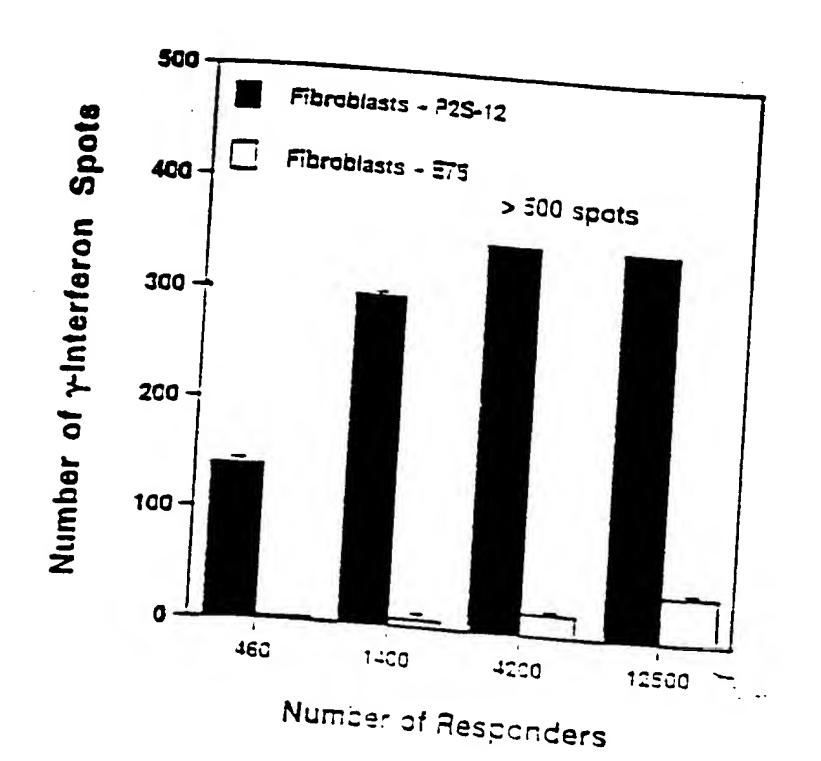


Fig. 2A

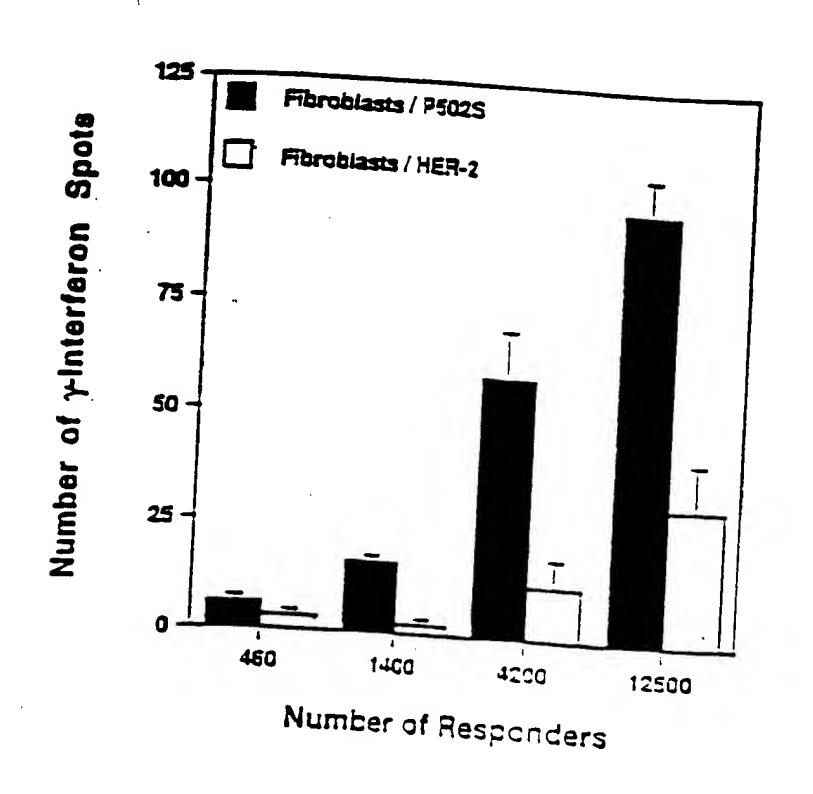


Fig. 2B

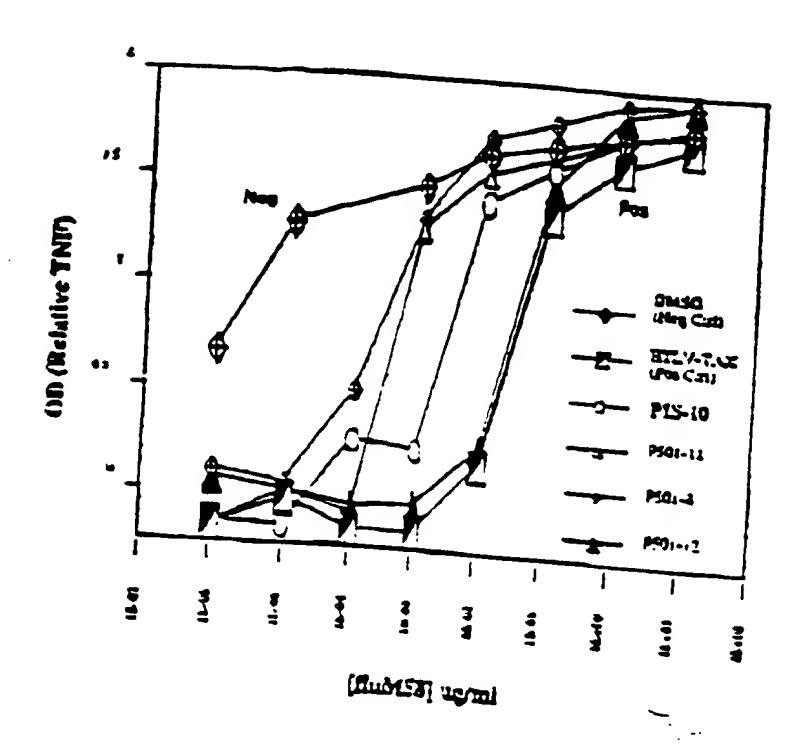


Fig. 3

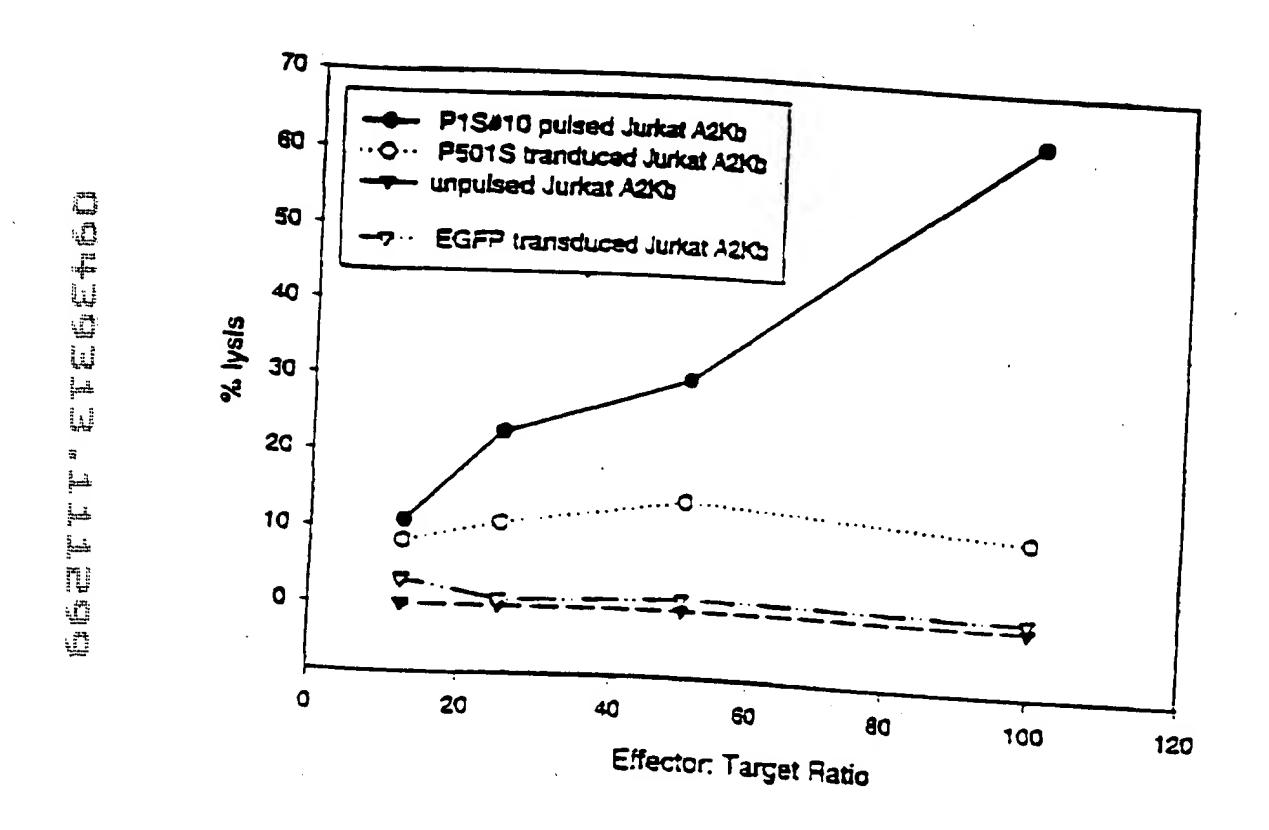


Fig. 4

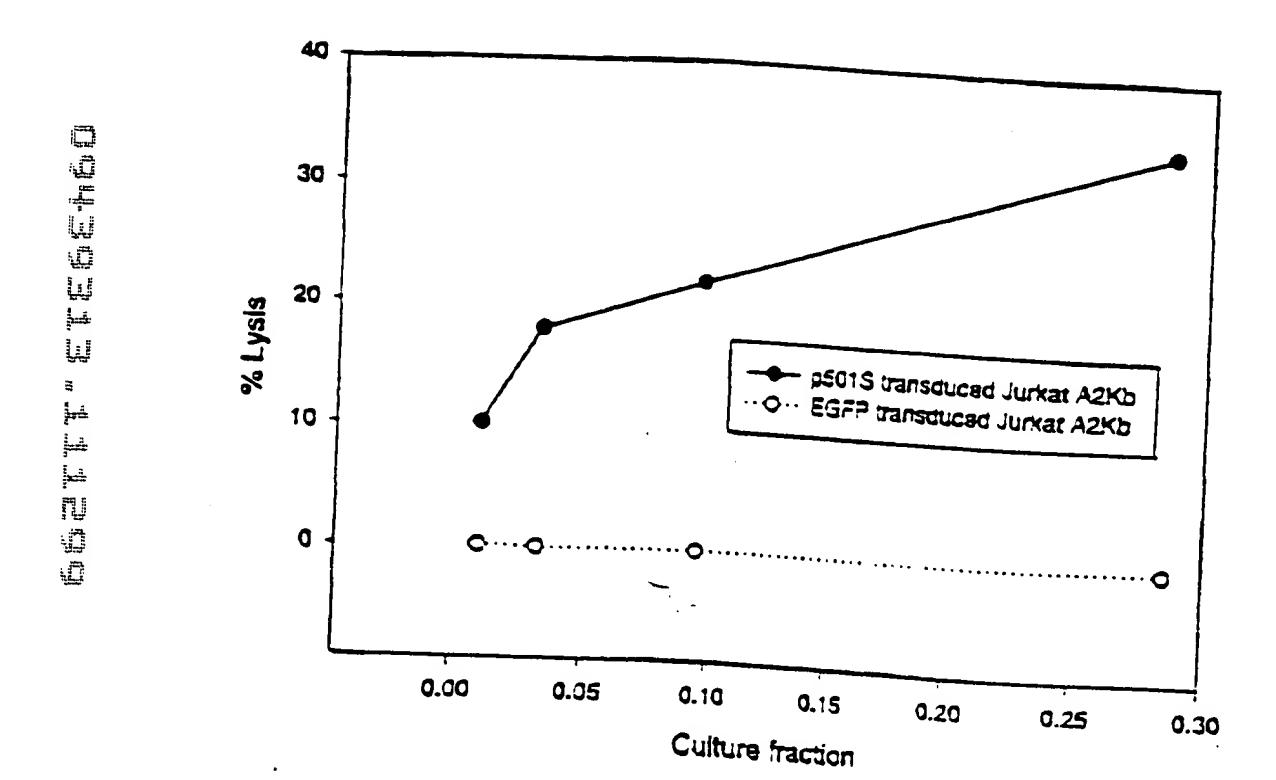
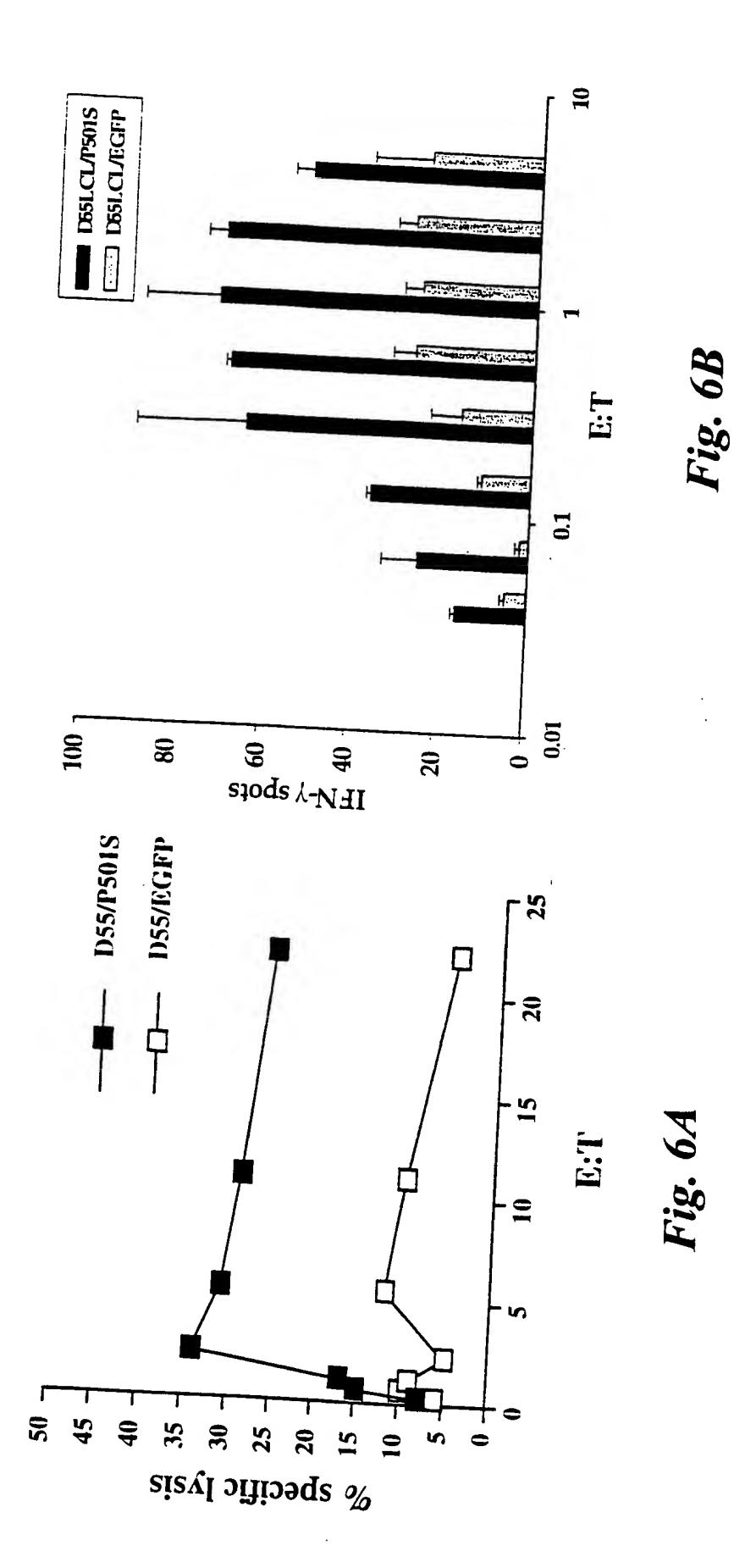
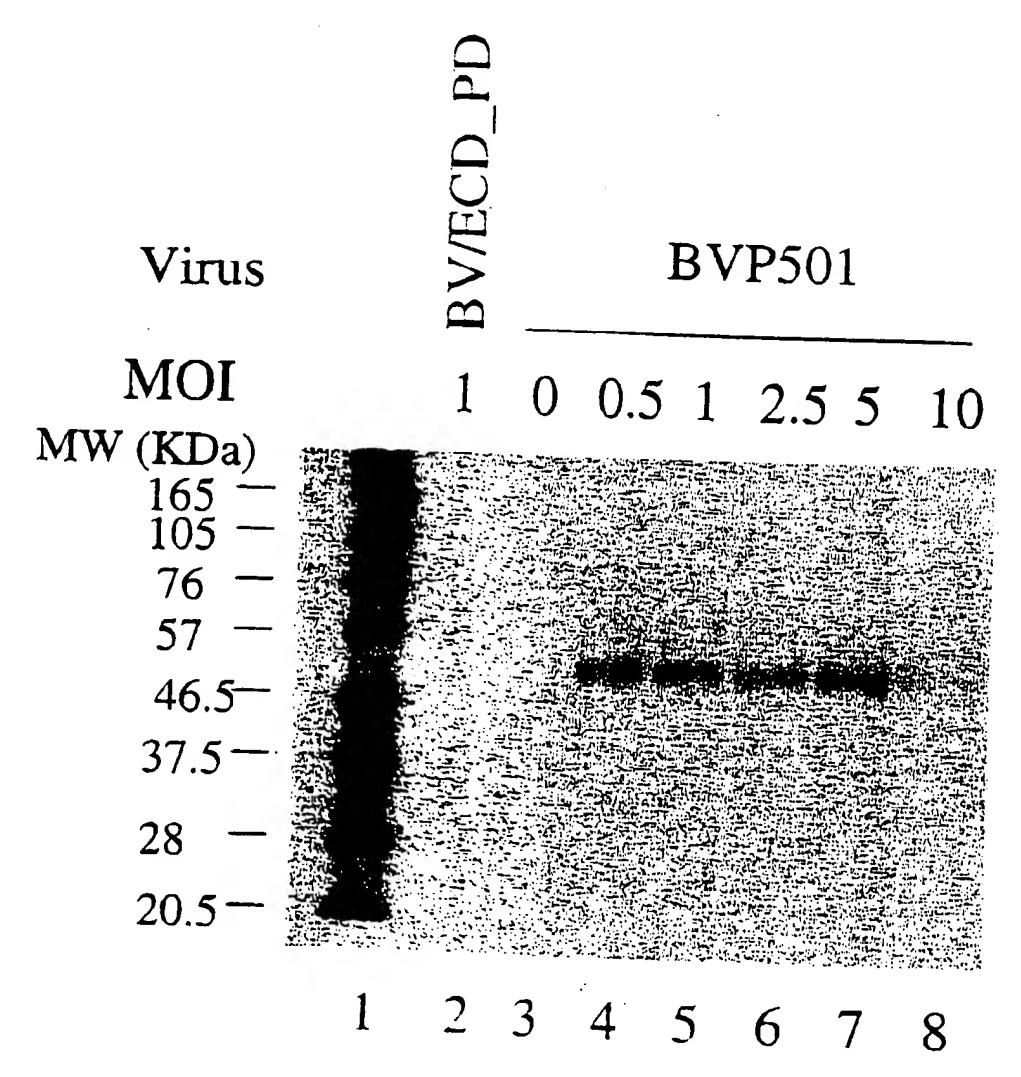


Fig. 5



## Expression of P501S by the Baculovirus Expression System



0.6 million high 5 cells in 6-well plate were infected with an unrelated control virus BV/ECD\_PD (lane 2), without virus (lane 3), or with recombinant baculovirus for P501 at different MOIs (lane 4 – 8). Cell lysates were run on SDS-PAGE under the reducing conditions and analyzed by Western blot with a monoclonal antibody against P501S (P501S-10E3-G4D3). Lane 1 is the biotinylated protein molecular weight marker (BioLabs).

Figure 8. Mapping of the epitope recognized by 10E3-G4-D3

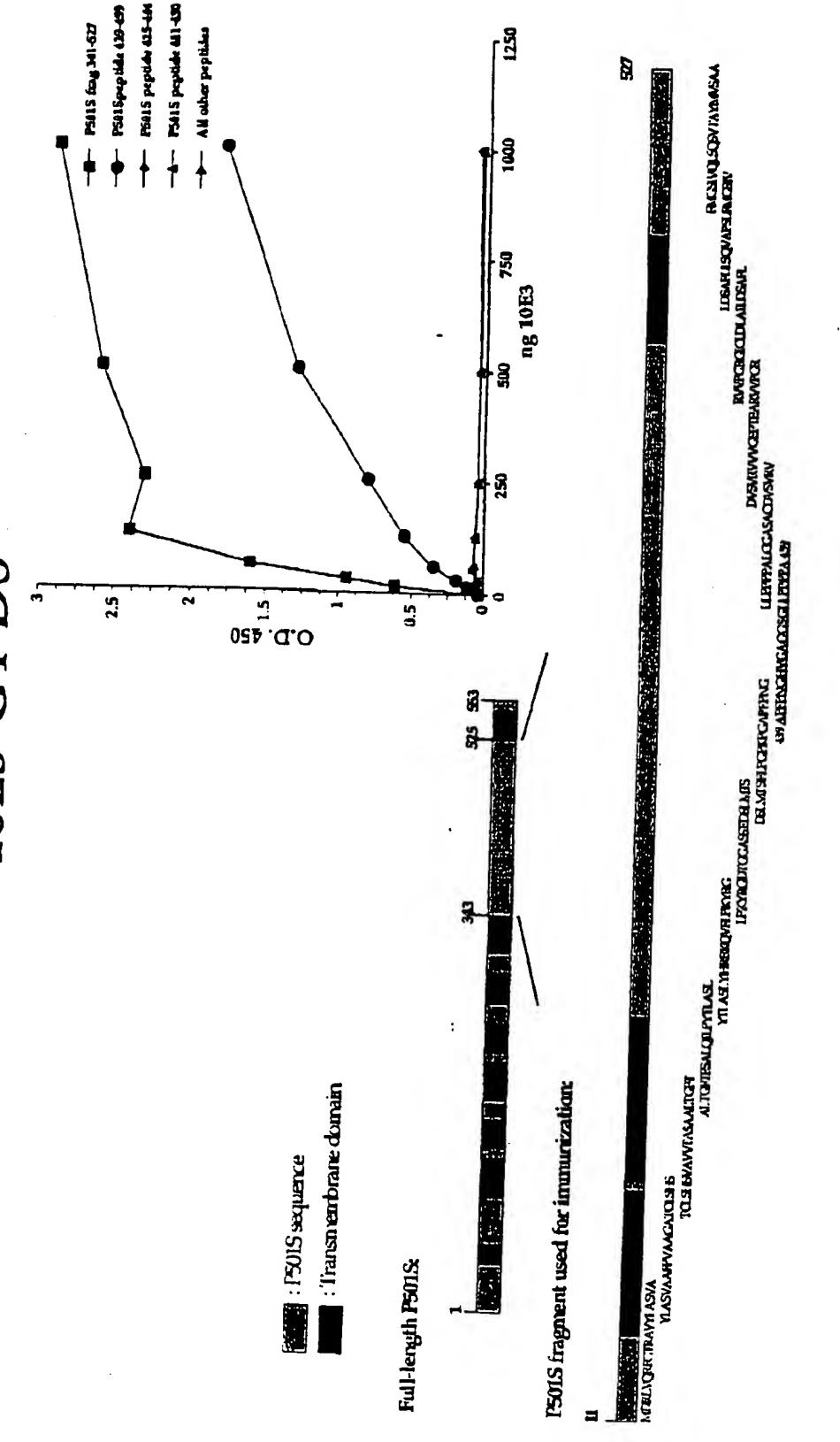


Fig. 8

## transmembrane, cytoplasmic, and extracellular regions Figure 1. Schematic of P501S with predicted

MVQRLWVSRLLRHRK AQLLLVNLLTFGLEVCLAAGIT YVPPI.LLEVGVEEKFM TMVLGIGPVLGLVCVPLLGSAS

DHWRGRYGRRP FIWALSLGILLSLFLIPRAGWL AGLLCPDPRPI.E LALL. II.GVGLLDFCGQVCFTPL

ISLGGCLGYLLPAL DWIYI'SALAPYLGTQEE EALLSULFRDPDHCRQ AYSVYAFM

CLFGLLTLIFI,TCVAATLLV AEEAALGPTEPAEGLSAPSI,SPHCCPCRARLAFRNLGALLPRL

HOLCCRMPRTLRR LFVAELCSWMALMTFTLFYTDF VGEGLYOGVPRARPGTEARRHYDEGVR

MGSLGLFLOCAISLVFSLVM DRLVQRFGTRAVYLAS VAAFPVAAGATCLSHSVAVVTA

LTGETFSALOIL PYTLASLY HREKQVFL PKYRGDTGGASSEDSLMTSFL PGPKPGAPFPNGIIVGAGGSGL

**MGSIVOLSOS** ICL DLAIL DSAFLL SOVAPSLF LPPPPALCGASACDVSVRVVVGEPTEARVVPGRG

## VTAYMVSAAGLGLVAIYFAT QVVFDKSDLAKYSA

Sequence in bold/underlined: used to generate polyclonal rabbit serum Underlined sequence: Predicted transmembrane domain; Bold sequence: Predicted extracellular domain; Italic sequence: Predicted intracellular domain.

Membrane Proteins: Applications to topology Prediction. J. Mul Biol. 283, OP (G.E. Tusnady and I. Simon (1998) Principles Localization of domains predicted using HMM1 Governing Amino Acid Composition of Integral 489-506.

Genomic Map of (5) Corixa Candidate Genes



Fig. 10

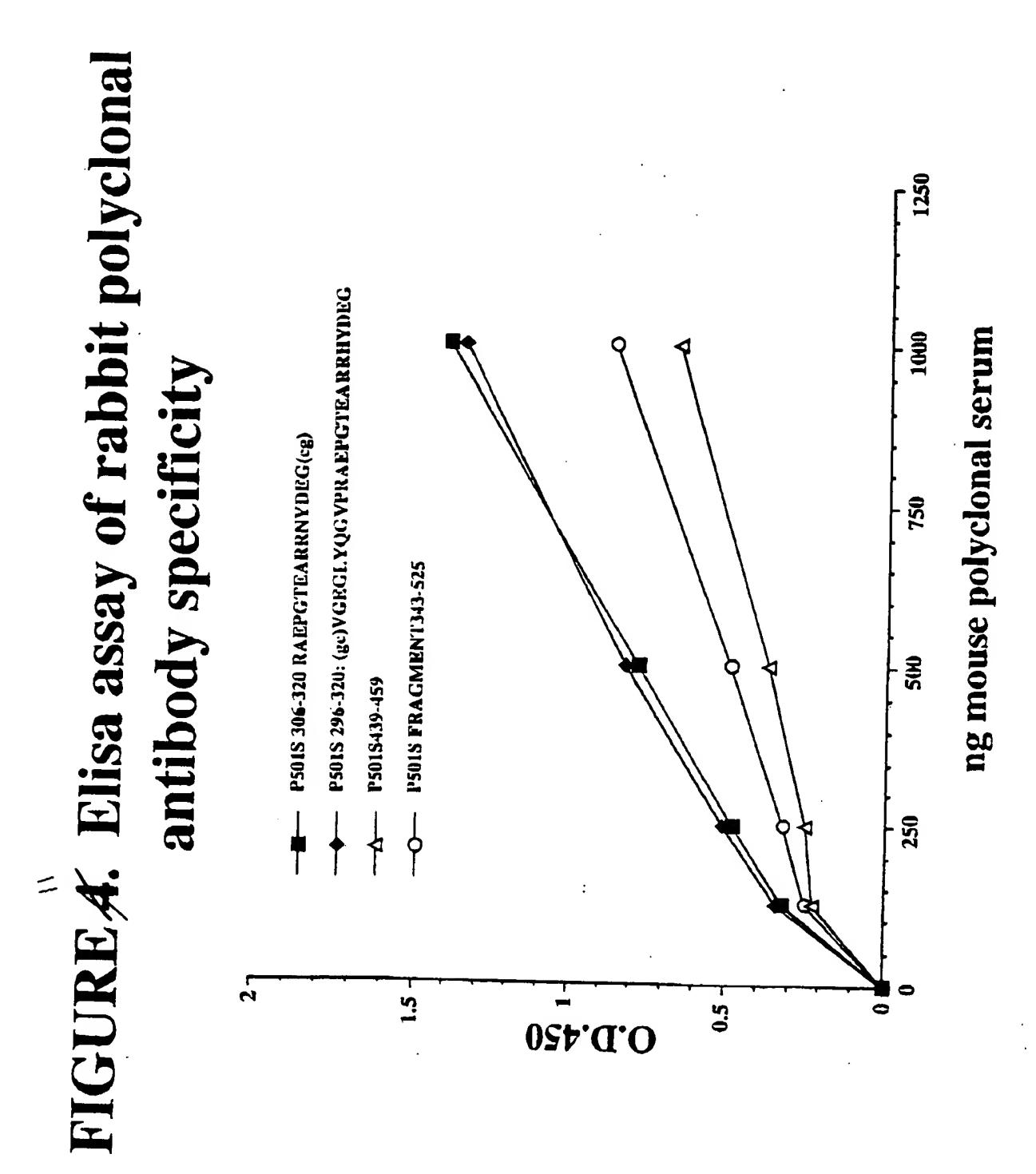


Fig. 11